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## RAW SEQUENCE LISTING

DATE: 02/25/2002

PATENT APPLICATION: US/10/050,726

TIME: 11:22:14

Input Set : N:\Crf3\RULE60\10050726.txt

Output Set: N:\CRF3\02252002\J050726.raw

5 <110> APPLICANT: Pathirana, Marie Sudam  
 9 <120> TITLE OF INVENTION: DNA ENCODING ORPHAN SNORF68 RECEPTOR  
 13 <130> FILE REFERENCE: 60795  
 15 <140> CURRENT APPLICATION NUMBER: 10/050,726  
 17 <141> CURRENT FILING DATE: 2002-01-16  
 19 <150> PRIOR APPLICATION NUMBER: 09/466,570  
 21 <151> PRIOR FILING DATE: 1999-12-17  
 25 <160> NUMBER OF SEQ ID NOS: 2  
 29 <170> SOFTWARE: PatentIn Ver. 2.0 - beta  
 33 <210> SEQ ID NO: 1  
 35 <211> LENGTH: 1536  
 37 <212> TYPE: DNA  
 39 <213> ORGANISM: Homo sapiens  
 43 <400> SEQUENCE: 1  
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 49 gacttctctc gcggccacgg cggccgtgct ctctctcagc accgtggcga ccgcggcgct 180  
 51 ggggaacctg agcgacgcaa gcggaggcgg cacagctgcc gctcccggtg gcggcggcct 240  
 53 tggcgggtcc ggggcagcgc gggaggcggg ggcgggcgtg aggcggccgc taggcccggg 300  
 55 ggcggcgccg ctgctgtcgc acggagctgc agtggcggcc caggcgctcg tctctctgct 360  
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 59 gcagctccgc accgtcacca acgccttcat cctgtcgtcg tccctatcgg atctgtctac 480  
 61 ggcgctgctc tgcttgcgg ccgccttctt gaacctcttc actccgcccc ggggttcggc 540  
 63 gcctgcggcc gccgcggggc cctggcgcg cttctgcgcc gccagccgct tcttcagctc 600  
 65 gtgcttcggc atcgtgtcca cgtcagcgt ggcgctcatc tcgttggacc gttactgcgc 660  
 67 tatcgtgcgg ccgcgcgggg agaagatcgg ccgccgccgc gcgctgcagc tgcgtggcgg 720  
 69 cgcttggttg acggccctgg gcttctctt gccctgggag ctgctcgggg cgccccggga 780  
 71 actcgcggcg gcgcagagct tccagggctg cctctaccgg acctccccg accccgcgca 840  
 73 gctgggcgcg gccttcagcg tggggctggg ggtggcctgc tacctgctgc ccttctctgt 900  
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 87 cgctttctct cccagccagg gcccggtct gcaagccaga agccgcagtc gccttcgaaa 1320  
 89 ccgctatgcc aaccggctgg gggcctgcaa caqatgtcc tcttccaacc cggccagcgg 1380  
 91 agtggcaggg gacgtggcca tgtgggcccc caaaaatcca gttgtacttt tctgcccaga 1440  
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 99 <210> SEQ ID NO: 2  
 101 <211> LENGTH: 494  
 103 <212> TYPE: PRT

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105 &lt;213&gt; ORGANISM: Homo sapiens

109 &lt;400&gt; SEQUENCE: 2

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111 Met Glu Glu Pro Gln Pro Pro Arg Pro Pro Ala Ser Met Ala Leu Leu
113   1           5           10           15
117 Gly Ser Gln His Ser Gly Ala Pro Ser Ala Ala Gly Pro Pro Gly Gly
119           20           25           30
123 Thr Ser Ser Ala Ala Thr Ala Ala Val Leu Ser Phe Ser Thr Val Ala
125           35           40           45
129 Thr Ala Ala Leu Gly Asn Leu Ser Asp Ala Ser Gly Gly Gly Thr Ala
131           50           55           60
135 Ala Ala Pro Gly Gly Gly Gly Leu Gly Gly Ser Gly Ala Ala Arg Glu
137   65           70           75           80
141 Ala Gly Ala Ala Val Arg Arg Pro Leu Gly Pro Glu Ala Ala Pro Leu
143           85           90           95
147 Leu Ser His Gly Ala Ala Val Ala Ala Gln Ala Leu Val Leu Leu Leu
149           100          105          110
153 Ile Phe Leu Leu Ser Ser Leu Gly Asn Cys Ala Val Met Gly Val Ile
155           115          120          125
159 Val Lys His Arg Gln Leu Arg Thr Val Thr Asn Ala Phe Ile Leu Ser
161           130          135          140
165 Leu Ser Leu Ser Asp Leu Leu Thr Ala Leu Leu Cys Leu Pro Ala Ala
167 145           150          155          160
171 Phe Leu Asp Leu Phe Thr Pro Pro Gly Gly Ser Ala Pro Ala Ala Ala
173           165          170          175
177 Ala Gly Pro Trp Arg Gly Phe Cys Ala Ala Ser Arg Phe Phe Ser Ser
179           180          185          190
183 Cys Phe Gly Ile Val Ser Thr Leu Ser Val Ala Leu Ile Ser Leu Asp
185           195          200          205
189 Arg Tyr Cys Ala Ile Val Arg Pro Pro Arg Glu Lys Ile Gly Arg Arg
191           210          215          220
195 Arg Ala Leu Gln Leu Leu Ala Gly Ala Trp Leu Thr Ala Leu Gly Phe
197 225           230          235          240
201 Ser Leu Pro Trp Glu Leu Leu Gly Ala Pro Arg Glu Leu Ala Ala Ala
203           245          250          255
207 Gln Ser Phe His Gly Cys Leu Tyr Arg Thr Ser Pro Asp Pro Ala Gln
209           260          265          270
213 Leu Gly Ala Ala Phe Ser Val Gly Leu Val Val Ala Cys Tyr Leu Leu
215           275          280          285
219 Pro Phe Leu Leu Met Cys Phe Cys His Tyr His Ile Cys Lys Thr Val
221           290          295          300
225 Arg Leu Ser Asp Val Arg Val Arg Pro Val Asn Thr Tyr Ala Arg Val
227 305           310          315          320
231 Leu Arg Phe Phe Ser Glu Val Arg Thr Ala Thr Thr Val Leu Ile Met
233           325          330          335
237 Ile Val Phe Val Ile Cys Cys Trp Gly Pro Tyr Cys Phe Leu Val Leu
239           340          345          350
243 Leu Ala Ala Ala Arg Gln Ala Gln Thr Met Gln Ala Pro Ser Leu Leu
245           355          360          365
249 Ser Val Val Ala Val Trp Leu Thr Trp Ala Asn Gly Ala Ile Asn Pro

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251	370	375	380
255	Val Ile Tyr Ala Ile Arg Asn Pro Asn Ile Ser Met Leu Leu Gly Arg		
257	385	390	395 400
261	Asn Arg Glu Glu Gly Tyr Arg Thr Arg Asn Val Asp Ala Phe Leu Pro		
263		405	410 415
267	Ser Gln Gly Pro Gly Leu Gln Ala Arg Ser Arg Ser Arg Leu Arg Asn		
269		420	425 430
273	Arg Tyr Ala Asn Arg Leu Gly Ala Cys Asn Arg Met Ser Ser Ser Asn		
275		435	440 445
279	Pro Ala Ser Gly Val Ala Gly Asp Val Ala Met Trp Ala Arg Lys Asn		
281		450	455 460
285	Pro Val Val Leu Phe Cys Arg Glu Gly Pro Pro Glu Pro Val Thr Ala		
287	465	470	475 480
291	Val Thr Lys Gln Pro Lys Ser Glu Ala Gly Asp Thr Ser Leu		
293		485	490

**VERIFICATION SUMMARY**

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